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OW protein - protein search, using sw model

Run on: September 4, 2002, 14:04:21 : Search time 99.82 Seconds  
(without alignments)  
114.763 Million cell updates/sec

Title: US-09-052-089A-1  
Perfect score: 2384  
Sequence: 1 MP1RALCTICSDPFDRHSRDV.....VRVKTVP5LPQAKIDFLWS 469

Scoring table: BIOSUM62  
Gapop 10.0 , Gapext 0.5

Searched: 231628 seqs, 24425594 residues

Total number of hits satisfying chosen parameters: 231628

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Issued\_Patents\_AA:\*  
1: /cgn2\_6/ptodata/2/1aa/5A.COMB.pep:\*  
2: /cgn2\_6/ptodata/2/1aa/5B.COMB.pep:\*  
3: /cgn2\_6/ptodata/2/1aa/6A.COMB.pep:\*  
4: /cgn2\_6/ptodata/2/1aa/6B.COMB.pep:\*  
5: /cgn2\_6/ptodata/2/1aa/PTCIS.COMB.pep:\*  
6: /cgn2\_6/ptodata/2/1aa/backfile1.pep:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	2384	100.0	469	4	US-09-052-089A-1
2	2357	98.9	469	2	US-08-968-751-2
3	1798.5	75.4	470	4	US-09-052-089A-2
4	1066	44.7	220	4	US-09-052-089A-3
5	906	38.0	220	4	US-09-052-089A-4
6	275	11.5	51	4	US-09-052-089A-5
7	272	11.4	51	4	US-09-052-089A-6
8	168.5	7.1	443	2	US-08-795-475-6
9	158	6.6	2482	1	US-08-328-254-6
10	158	6.6	3248	1	US-08-353-700-1
11	157	6.6	3248	5	PCT-US95-16216-1
12	157	6.6	756	4	US-09-085-1199B-9
13	156	6.5	976	4	US-09-104-324B-4
14	156	6.5	1939	4	US-09-310-187A-1
15	154	6.5	1886	4	US-08-938-105-3
16	152	6.4	2101	1	US-08-466-390-4
17	152	6.4	2101	1	US-08-470-950-4
18	152	6.4	2101	1	US-08-467-781-4
19	152	6.4	2101	1	US-08-195-487-4
20	152	6.4	2101	2	US-08-483-924-4
21	152	6.4	2101	5	US-09-452-294-1
22	149	6.2	896	1	US-08-095-737-2
23	149	6.2	896	1	US-08-480-145-2
24	149	6.2	896	2	US-08-477-389-2
25	149	6.2	896	2	US-08-477-389-2
26	149	6.2	1093	5	PCT-US93-03077-1
27	148.5	6.2	1312	2	US-08-687-080-51

28	147.5	6.2	316	4	US-08-098-327E-31	Sequence 31, Appl
29	147.5	6.2	316	4	US-08-462-625-31	Sequence 31, Appl
30	147.5	6.2	1090	4	US-09-085-199B-5	Sequence 5, Appl
31	147.5	6.2	1312	2	US-08-592-126-148	Sequence 148, App
32	146.5	6.1	914	4	US-09-085-199B-4	Sequence 4, Appl
33	146	6.1	1068	4	US-09-085-199B-11	Sequence 11, Appl
34	145.5	6.1	1713	3	US-08-600-982-24	Sequence 24, Appl
35	145.5	6.1	1713	3	PCT-US94-10261A-24	Sequence 24, Appl
36	144	6.0	376	6	5180810-1	Patent No. 5180810
37	144	6.0	414	5	PCT-US93-03077-3	Sequence 3, Appl
38	142.5	6.0	816	2	US-08-533-306A-6	Sequence 6, Appl
39	142.5	6.0	816	2	US-08-742-923A-6	Sequence 6, Appl
40	142.5	6.0	885	2	US-08-533-306A-4	Sequence 4, Appl
41	142.5	6.0	885	2	US-08-742-923A-4	Sequence 4, Appl
42	140	5.9	576	2	US-08-533-306A-2	Sequence 2, Appl
43	140	5.9	576	2	US-08-742-923A-2	Sequence 2, Appl
44	139.5	5.9	1360	4	US-09-393-569-2	Sequence 2, Appl
45	139	5.8	386	4	US-09-085-199B-2	Sequence 2, Appl

ALIGNMENTS

RESULT 1  
US-09-052-089A-1  
Sequence 1, Application US/09052089A  
Patent No. 6346505  
GENERAL INFORMATION:  
APPLICANT: Lee, Soo Y.  
TITLE OF INVENTION: ChOI, Yongwon  
FAMILY, AND USES THEREOF  
NUMBER OF SEQUENCES: 16  
CORRESPONDENCE ADDRESSES:  
ADDRESSEE: David A. Jackson, Esq.  
STREET: 411 Hackensack Ave, Continental Plaza, 4th Floor  
CITY: Hackensack  
STATE: New Jersey  
COUNTRY: USA  
ZIP: 07601  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/052,089A  
FILING DATE: 31-Mar-1998  
CLASSIFICATION: <Unknown>  
ATTORNEY/AGENT INFORMATION:  
NAME: Jackson Esq., David A.  
REGISTRATION NUMBER: 26,742  
REFERENCE/DOCKET NUMBER: 600-1-198 CIP 1  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 201-487-5800  
TELEFAX: 201-343-1684  
INFORMATION FOR SEQ ID NO: 1:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 469 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
HYPOTHETICAL: NO  
FRAGMENT TYPE: <Unknown>  
ORIGINAL SOURCE:  
ORGANISM: Homo sapiens  
SEQUENCE DESCRIPTION: SEQ ID NO: 1:  
US-09-052-089A-1  
Query Match 100.0%; Score 2384; DB 4; Length 469;

Best Local Similarity 100.0%; Pred. No. 1.1e-196;  
Matches 469; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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QY 1 MPRLACTTCSDFPDHSDVAIAHCGHTFHLQCLIOSFETAPSRTPCQCRIOVGKRTIIN 60
Db 1 MPRLACTTCSDFPDHSDVAIAHCGHTFHLQCLIOSFETAPSRTPCQCRIOVGKRTIIN 60
QY 61 KLFEDLAQEEENVLDREFLNKNELDNVRAQLSQDKERKRSQVYIIDTLRDTLEERNATVVS 120
Db 61 KLFEDLAQEEENVLDREFLNKNELDNVRAQLSQDKERKRSQVYIIDTLRDTLEERNATVVS 120
QY 121 LQALGKAEMLCSTLKQKMYLEQOODETKQAQEEAGRLRSKMKTMQEIELLQSQLPYV 180
Db 121 LQALGKAEMLCSTLKQKMYLEQOODETKQAQEEAGRLRSKMKTMQEIELLQSQLPYV 180
QY 181 EEMIRDMGVGQSAVEOLAVYCVSLKKEYENLKEARKASGEVADKLKRDLFSSRSKLQTVY 240
Db 181 EEMIRDMGVGQSAVEOLAVYCVSLKKEYENLKEARKASGEVADKLKRDLFSSRSKLQTVY 240
QY 241 SELDQAKLELKSQKDLQSDADKEIMSLKRLTLMQETLNLPPVASETVDRVLYESPAVE 300
Db 241 SELDQAKLELKSQKDLQSDADKEIMSLKRLTLMQETLNLPPVASETVDRVLYESPAVE 300
QY 301 VNLKLRPSFRDDIDLNATFDVDTPPARPSSQHGYYEKLCLKESHSPIQDVPRKTKCKGP 360
Db 301 VNLKLRPSFRDDIDLNATFDVDTPPARPSSQHGYYEKLCLKESHSPIQDVPRKTKCKGP 360
QY 361 RKESQSLSGQSCAGPDEDELVGAFPIFVRNALLGOKPRPSSSCSKDVYRTGFDGL 420
Db 361 RKESQSLSGQSCAGPDEDELVGAFPIFVRNALLGOKPRPSSSCSKDVYRTGFDGL 420
QY 421 GGRTKFIQPTDVMIRPLPVKPKTKVKQRYRVTVPSLFOAKLDTFLMS 469
Db 421 GGRTKFIQPTDVMIRPLPVKPKTKVKQRYRVTVPSLFOAKLDTFLMS 469

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RESULT 2  
US-08-968-751-2  
Sequence 2, Application US/08968751  
Patent No. 5948643

GENERAL INFORMATION:  
APPLICANT: Rudinfeld, Bonnee  
APPLICANT: Polakis, Paul G.  
APPLICANT: Ligentel, Carol  
APPLICANT: Vuong, Terilyn T.  
TITLE OF INVENTION: MODULATORS OF BRCA1 ACTIVITY  
NUMBER OF SEQUENCES: 6  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: ONYX Pharmaceuticals, Inc.  
STREET: 3031 Research Drive  
CITY: Richmond  
STATE: CA  
COUNTRY: USA  
ZIP: 94806

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/968,751  
FILING DATE:  
CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:  
NAME: Giotta, Gregory  
REGISTRATION NUMBER: 32,028  
REFERENCE/DOCKET NUMBER: ONYX1024 GG  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (510) 262-8710  
TELEFAX: (510) 222-9758  
INFORMATION FOR SEQ ID NO: 2:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 469 amino acids

TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-968-751-2

Query Match  
Best Local Similarity 98.9%; Score 2357; DB 2; Length 469;  
Matches 465; Conservative 0; Mismatches 4; Indels 0; Gaps 0;

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QY 1 MPRLACTTCSDFPDHSDVAIAHCGHTFHLQCLIOSFETAPSRTPCQCRIOVGKRTIIN 60
Db 1 MPRLACTTCSDFPDHSDVAIAHCGHTFHLQCLIOSFETAPSRTPCQCRIOVGKRTIIN 60
QY 61 KLFEDLAQEEENVLDREFLNKNELDNVRAQLSQDKERKRSQVYIIDTLRDTLEERNATVVS 120
Db 61 KLFEDLAQEEENVLDREFLNKNELDNVRAQLSQDKERKRSQVYIIDTLRDTLEERNATVVS 120
QY 121 LQALGKAEMLCSTLKQKMYLEQOODETKQAQEEAGRLRSKMKTMQEIELLQSQLPYV 180
Db 121 LQALGKAEMLCSTLKQKMYLEQOODETKQAQEEAGRLRSKMKTMQEIELLQSQRPYV 180
QY 181 EEMIRDMGVGQSAVEOLAVYCVSLKKEYENLKEARKASGEVADKLKRDLFSSRSKLQTVY 240
Db 181 EEMIRDMGVGQSAVEOLAVYCVSLKKEYENLKEARKASGEVADKLKRDLFSSRSKLQTVY 240
QY 241 SELDQAKLELKSQKDLQSDADKEIMSLKRLTLMQETLNLPPVASETVDRVLYESPAVE 300
Db 241 SELDQAKLELKSQKDLQSDADKEIMSLKRLTLMQETLNLPPVASETVDRVLYESPAVE 300
QY 301 VNLKLRPSFRDDIDLNATFDVDTPPARPSSQHGYYEKLCLKESHSPIQDVPRKTKCKGP 360
Db 301 VNLKLRPSFRDDIDLNATFDVDTPPARPSSQHGYYEKLCLKESHSPIQDVPRKTKCKGP 360
QY 361 RKESQSLSGQSCAGPDEDELVGAFPIFVRNALLGOKPRPSSSCSKDVYRTGFDGL 420
Db 361 RKESQSLSGQSCAGPDEDELVGAFPIFVRNALLGOKPRPSSSCSKDVYRTGFDGL 420
QY 421 GGRTKFIQPTDVMIRPLPVKPKTKVKQRYRVTVPSLFOAKLDTFLMS 469
Db 421 GGRTKFIQPTDVMIRPLPVKPKTKVKQRYRVTVPSLFOAKLDTFLMS 469

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RESULT 3  
US-09-052-089a-2  
Sequence 2, Application US/09052089A  
Patent No. 6346605

GENERAL INFORMATION:  
APPLICANT: Lee, Soo Y.  
APPLICANT: Choi, Yongwon  
TITLE OF INVENTION: SIGNAL TRANSDUCER FOR THE TNF RECEPTOR SUPER  
FAMILY, AND USES THEREOF  
NUMBER OF SEQUENCES: 16  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: David A. Jackson, Esq.  
STREET: 411 Hackensack Ave, Continental Plaza, 4th  
FLOOR  
CITY: Hackensack  
STATE: New Jersey  
COUNTRY: USA  
ZIP: 07601

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/052,089A  
FILING DATE: 31-Mar-1998  
CLASSIFICATION: <Unknown>

ATTORNEY/AGENT INFORMATION:  
NAME: Jackson Esq., David A.  
REGISTRATION NUMBER: 26,742



COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/052,089A  
FILING DATE: 31-Mar-1998  
CLASSIFICATION: <Unknown>  
ATTORNEY/AGENT INFORMATION:  
NAME: Jackson Esq., David A.  
REGISTRATION NUMBER: 26,742  
REFERENCE/DOCKET NUMBER: 600-1-198 CIP 1  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 201-487-5800  
TELEFAX: 201-343-1684  
INFORMATION FOR SEQ ID NO: 4:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 220 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
FRAGMENT TYPE: Internal  
ORIGINAL SOURCE:  
ORGANISM: mouse  
SEQUENCE DESCRIPTION: SEQ ID NO: 4:  
US-09-052-089A-4

Query Match 38.0%; Score 906; DB 4; Length 220;  
Best Local Similarity 86.0%; Pred. No. 2.8e-70;  
Matches 185; Conservative 18; Mismatches 12; Indels 0; Gaps 0;

QY 56 RTIINKLFFDLAEEENVLDREFLKNELDNVRALDSQDKKRSQYIITLRLTEERN 115  
DB 1 KTIINKLFFDLAEEENVLDREFLKNELDNVRALDSQDKKRSQYIITLRLTEERN 60  
QY 116 ATVVSLOALGKAMLCSTLKKOKKYLEQODEFKQAQOEGRLSKMKTKTEQJELLLOS 175  
DB 61 ATVVSLOALGKAMLCSTLKKOKKYLEQODEFKQAQOEGRLSKMKTKTEQJELLLOS 120  
QY 176 QLPVEEMIRDMGVGSAVEQLAVYCVSLKKEYENLKEARRASGEVADKLKDLFSSRSK 235  
DB 121 QLPVEEMIRDMGVGSAVEQLAVYCVSLKKEYENLKEARRASGEVADKLKDLFSSRSK 180  
QY 236 LQTYSELDAQKLELSAOKDLOSADKEIMSLRKK 270  
DB 181 LKTLNTELDQAKLELSAOKDLOSADKEIMSLRKK 215

RESULT 6  
US-09-052-089A-5  
Sequence 5, Application US/09052089A  
Patent No. 6346605  
GENERAL INFORMATION:  
APPLICANT: Lee, Soo Y.  
Choi, Yongwon  
TITLE OF INVENTION: SIGNAL TRANSDUCER FOR THE TNP RECEPTOR SUPER  
FAMILY, AND USES THEREOF  
NUMBER OF SEQUENCES: 16  
CORRESPONDENCE ADDRESSES:  
ADDRESSEE: David A. Jackson, Esq.  
STREET: 411 Hackensack Ave, Continental Plaza, 4th  
Floor  
CITY: Hackensack  
STATE: New Jersey  
COUNTRY: USA  
ZIP: 07601  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/052,089A  
FILING DATE: 31-Mar-1998  
CLASSIFICATION: <Unknown>  
ATTORNEY/AGENT INFORMATION:  
NAME: Jackson Esq., David A.  
REGISTRATION NUMBER: 26,742  
REFERENCE/DOCKET NUMBER: 600-1-198 CIP 1  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 201-487-5800  
TELEFAX: 201-343-1684  
INFORMATION FOR SEQ ID NO: 5:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 51 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
HYPOTHETICAL: NO  
FRAGMENT TYPE: Internal  
ORIGINAL SOURCE:  
ORGANISM: Homo sapiens  
SEQUENCE DESCRIPTION: SEQ ID NO: 5:  
US-09-052-089A-5

Query Match 11.5%; Score 275; DB 4; Length 51;  
Best Local Similarity 96.1%; Pred. No. 5.2e-17;  
Matches 49; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 4 RALCTICSDFFDHSRDVAALHCHTFHLQCLIOSFETAPSRTPCPOCRIOYG 54  
DB 1 RALCTICSDFFDHSRDVAAMDCGHTFHLQCLIOSFETAPSRTPCPOCRIOYG 51

RESULT 7  
US-09-052-089A-6  
Sequence 6, Application US/09052089A  
Patent No. 6346605  
GENERAL INFORMATION:  
APPLICANT: Lee, Soo Y.  
Choi, Yongwon  
TITLE OF INVENTION: SIGNAL TRANSDUCER FOR THE TNP RECEPTOR SUPER  
FAMILY, AND USES THEREOF  
NUMBER OF SEQUENCES: 16  
CORRESPONDENCE ADDRESSES:  
ADDRESSEE: David A. Jackson, Esq.  
STREET: 411 Hackensack Ave, Continental Plaza, 4th  
Floor  
CITY: Hackensack  
STATE: New Jersey  
COUNTRY: USA  
ZIP: 07601  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patentin Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/052,089A  
FILING DATE: 31-Mar-1998  
CLASSIFICATION: <Unknown>  
ATTORNEY/AGENT INFORMATION:  
NAME: Jackson Esq., David A.  
REGISTRATION NUMBER: 26,742  
REFERENCE/DOCKET NUMBER: 600-1-198 CIP 1  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 201-487-5800  
TELEFAX: 201-343-1684  
INFORMATION FOR SEQ ID NO: 6:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 51 amino acids  
TYPE: amino acid

STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: peptide  
HYPOTHETICAL: NO  
FRAGMENT TYPE: internal  
ORIGINAL SOURCE:  
ORGANISM: mouse  
SEQUENCE DESCRIPTION: SEQ ID NO: 6  
US-09-052-089a-6

Query Match 11.4%; Score 272; DB 4; Length 51;  
Best Local Similarity 96.0%; Pred. No. 9,4e-17;  
Matches 48; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 5 ALCCTGSDFFDHSRDVAATHCGHTFHLCLOIOSEFAPSTCTCQCRIQVG 54  
DB 2 SLCTTCSDFDHSRDVAATHCGHTFHLCLOIOSEFAPSTCTCQCRIQVG 51

## RESULT 8

US-08-795-475-6  
Sequence 6, Application US/08795475  
Patent No. 5965390  
GENERAL INFORMATION:  
APPLICANT: Bjvrck, Lars  
TITLE OF INVENTION: PROTEIN L AND HYBRID PROTEINS THEREOF  
NUMBER OF SEQUENCES: 14  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: SEED AND BERRY LLP  
STREET: 6300 Columbia Center, 701 Fifth Avenue  
CITY: Seattle  
STATE: Washington  
COUNTRY: USA  
ZIP: 98104-7092  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
FILING DATE: 11-FEB-1997  
APPLICATION NUMBER: US/08/795,475  
CLASSIFICATION: 435  
ATTORNEY/AGENT INFORMATION:  
NAME: Mcmasters, David D.  
REGISTRATION NUMBER: 33,963  
REFERENCE/DOCKET NUMBER: 100084,402D1  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (206) 622-4900  
TELEFAX: (206) 682-6031  
INFORMATION FOR SEQ ID NO: 6:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 443 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-795-475-6

Query Match 7.1%; Score 168.5; DB 2; Length 443;  
Best Local Similarity 20.7%; Pred. No. 1.5e-06;  
Matches 73; Conservative 76; Mismatches 136; Indels 67; Gaps 11;

QY 64 FDLAQEEENLVDEFLKNELDNVRAQLSQK-DREKRDQVYIIDLRDTLEERNATVVSQ 122  
DB 78 YDLAKESTS-WDRQLEKLEKEKKEALEIDAQASRDYHATALEKELEKKKALELAI 136  
QY 123 QALGKRAMCSTLKKQKMYLEQODETK---QAQAEAGLRKSKM--TMEQLELLSQ 176  
DB 137 QA-SQDYNRANVALEKETTTREQEIINRLNLGNAKLELDQLSSKEQLTEKAKLEBEKQ 195

QY 177 LPEV--EEMIRMGVQSAVEQALVAVYCSLKKREYENLKEARRASGEVADLKRDLFFSSR- 233  
DB 196 ISDASRQSLRDLDDASREAKKQVEKDLNLTAEIDKVKDKQSDASRQLRDLDDASRE 255  
QY 234 -----SKLQTYSELDQAKLE-----LKSQKDLQASADEINS 266  
DB 256 AKQVEKDLNLTAEIDKVEEKQISDASRQLRDLDDASREAKKQVEKALEBANSKLAA 315  
QY 267 LKKKLTMLQETLNLPPVASETVDRVLLESPAPVEVNLKLRPSFRDIDLNA----- 318  
DB 316 LEKLNKELES-----KLTKEKAEIQAKLEAPAKLQKQLEAKLQKQLEAKLRA 364  
QY 319 --TFVDTPPARPSSQHGVEKLEKSHSPIQDVPKKICKGPRRESQSL 368  
DB 365 GKASDSQTFDTKPGN-----KAVPGKQAPQAGIKRPQNNKAPMKETKROL 409

## RESULT 9

US-08-328-254-6  
Sequence 6, Application US/08328254  
Patent No. 5710022  
GENERAL INFORMATION:  
APPLICANT: Zhu, Xueliang  
APPLICANT: Lee, Wen-Hwa  
TITLE OF INVENTION: A No. 5710022e1 Nuclear Mitotic Phosphoprotein  
NUMBER OF SEQUENCES: 8  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Campbell and Flores  
STREET: 4370 La Jolla Village Drive, Suite 700  
CITY: San Diego  
STATE: California  
COUNTRY: USA  
ZIP: 92122  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
FILING DATE: 24-OCT-1994  
APPLICATION NUMBER: US/08/328,254  
CLASSIFICATION: 435  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US 08/141,239  
FILING DATE: 22-OCT-1993  
ATTORNEY/AGENT INFORMATION:  
NAME: Campbell, Cathryn A.  
REGISTRATION NUMBER: 31,815  
REFERENCE/DOCKET NUMBER: P-CJ 1191  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (619) 535-9001  
TELEFAX: (619) 535-8949  
INFORMATION FOR SEQ ID NO: 6:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 2482 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-328-254-6

Query Match 6.6%; Score 158; DB 1; Length 2482;  
Best Local Similarity 23.2%; Pred. No. 0.00014;  
Matches 56; Conservative 55; Mismatches 94; Indels 36; Gaps 6;

QY 74 LDREFLKNELDNVRAQLSQKDEKRDQVYIIDLRDTLEERNATVVSQALQALKA-EMLC 132  
DB 1571 LDVLTLRSEKENLTQKQKQGLSELDKLSSFKSLBEKQAEIQIKRESSTAVEMQ 1630  
QY 133 STLAKQ-----WKYLEQOOD---ETKQAEAGLRKSKMTMEQLELLSQ 175  
DB 1631 NQKELNEAVALAQDQEIWKATEQSIDPPIEEHOLRNSIKELRLARLEADEKQKOLCVLD 1690

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0Y 176 QLP-----VEEMIDMGVGSANEQIAYVCSLKKREYENLKARAKASEVADK 224
Db 1691 QLKSEHHADLLKGRVENLEIRELEIARTOEHAALEANSKEVEFLNAKITGEMTQSLRG 1750
0Y 225 LRKDLFSSRSLQTVYSELQ-----AKLEL--KSAOKDLOSADKEIMSLKKLTMLOET 277
Db 1751 LELDVVITRSEKENLEIQLQEOERISLSELIINSSFENITLOEKDEOKVOMKEKSSTAME 1810
0Y 278 L 278
Db 1811 L 1811

```

RESULT 10  
 US-08-353-700-1  
 Sequence 1, Application US/08353700  
 Patent No. 5599919  
 GENERAL INFORMATION:  
 APPLICANT: YEN, TIMOTHY J.  
 APPLICANT: RATNER, JEROME B.  
 TITLE OF INVENTION: NUCLEIC ACID ENCODING A  
 TITLE OF INVENTION: TRANSIENTLY-EXPRESSED KINTOCHORE PROTEIN  
 TITLE OF INVENTION: AND METHODS OF USE  
 NUMBER OF SEQUENCES: 4  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: DANN, DOFFMAN, HERRELL AND SKILLMAN  
 STREET: 1601 MARKET STREET, SUITE 720  
 CITY: PHILADELPHIA  
 STATE: PA  
 COUNTRY: USA  
 ZIP: 19103-2307  
 COMPUTER READABLE FORM:  
 MEDIUM TYPE: floppy disk  
 COMPUTER: IBM PC compatible  
 OPERATING SYSTEM: PC-DOS/MS-DOS  
 SOFTWARE: PatentIn Release #1.0, Version #1.25  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/08/353,700  
 FILING DATE: 09-DEC-1994  
 CLASSIFICATION: 435  
 ATTORNEY/AGENT INFORMATION:  
 NAME: REED, JANET E.  
 REGISTRATION NUMBER: 36,252  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: (215) 563-4100  
 TELEFAX: (215) 563-4044  
 INFORMATION FOR SEQ ID NO: 1:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 3248 amino acids  
 TYPE: amino acid  
 STRANDEDNESS: single  
 TOPOLOGY: linear  
 MOLECULE TYPE: protein  
 HYPOTHETICAL: NO  
 ANTI-SENSE: NO  
 ORIGINAL SOURCE:  
 ORGANISM: HUMAN  
 US-08-353-700-1

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Query Match      6.6% Score 158 DB 1 Length 3248;
Best Local Similarity 23.2% Pred No. 0.00021;
Matches 56; Conservative 55; Mismatches 94; Indels 36; Gaps 6

QY    74 LDREFLNELNVRNQSLQSKDEKRDGYITLRTTEERNATVVSLOALGKA-EMLC 132
      ||::|::|::|::|::|::|::|::|::|::|::|::|::|::|::|::|
Db    2299 LDIYLRSEKENLTQIQIEKQQLSELKLSSPFSLEEKDAEQTQIEESTAVEMQ 2358

QY    133 STLKQ-----MKYLEOOD---ETKQAQEAAGRLRSKMTEQIEDLLQS 175
      ::||::|::|::|::|::|::|::|::|::|::|::|::|::|::|::|
Db    2359 NQLKELNEVAVALCGDQIEIMKTAEGLSDPIIEEHQILRISIKRLRLADEKKQCIVQ 2418

QY    176 QLPF-----VEEMIRDMGVGSAAVEDLAYVCVSLKEYEVLKFAARASGEVADK 224
```

Db	2419	QKRESEHHADLLGRVENELERELEIRTOGEHAALAEKSEKVEVETLAKKIEGMOJSLRG	2478
QY	225	LKRDDESRSKKLOTVYSELDQ---AKLEL--KSAQKDLOSGADKEIMSLKKKLTMLQET	277
Db	2479	LELDVYVIRSEKENLTELQEOERISELEIINSSFEINTELQEOEKVOMKEKSSTAMEM	2538
QY	278	L	278
Db	2539	L	2539

RESULT 11  
 PCT-US95-16216-1  
 : Sequence 1, Application PC/TUS9516216  
 : GENERAL INFORMATION:  
 : APPLICANT: Yen, Timothy J.  
 : APPLICANT: Rattner, Jerome B.  
 : TITLE OF INVENTION: Nucleic Acid Encoding a Transiently  
 : TITLE OF INVENTION: Expressed Kinetochore Protein, and Methods of Use  
 : NUMBER OF SEQUENCES: 4  
 : CORRESPONDENCE ADDRESS:  
 : ADDRESSEE: Dann, Dorfman, Herrell and Skillman  
 : STREET: 1601 Market Street Suite 720  
 : CITY: Philadelphia  
 : STATE: PA  
 : COUNTRY: USA  
 : ZIP: 19103-2307  
 : COMPUTER READABLE FORM:  
 : MEDIUM TYPE: Floppy disk  
 : COMPUTER: IBM PC compatible  
 : OPERATING SYSTEM: PC-DOS/MS-DOS  
 : SOFTWARE: PatentIn Release #1.0, Version #1.30  
 : CURRENT APPLICATION DATA:  
 : APPLICATION NUMBER: PCT/US95/16216  
 : FILING DATE:  
 : CLASSIFICATION:  
 : PRIOR APPLICATION DATA:  
 : APPLICATION NUMBER: US 08/353,700  
 : FILING DATE: 09-DEC-1995  
 : ATTORNEY/AGENT INFORMATION:  
 : NAME: Reed, Janet E.  
 : REGISTRATION NUMBER: 36,252  
 : TELECOMMUNICATION INFORMATION:  
 : TELEPHONE: (215) 563-4100  
 : TELEFAX: (215) 563-4004  
 : INFORMATION FOR SEQ ID NO: 1:  
 : SEQUENCE CHARACTERISTICS:  
 : LENGTH: 3248 amino acids  
 : TYPE: amino acid  
 : STRANDEDNESS: not relevant  
 : TOPOLOGY: not relevant  
 : MOLECULE TYPE: protein  
 : HYPOTHEICAL: NO  
 : ANTI-SENSE: NO  
 : PCT-US95-16216-1

[illegible]



Db 590 VKCKLSESENCNNLRKOVENKNRYIEBLOENK-ALKKGTAEKOLNVEIKVKNKLEL 648  
QY 172 LLOSO-----LPEVE--EMIRDMGVG-----QSAVE 195  
Db 649 ELBSAKOKFGFITDTYOKETIEDKRISEENLLEVEKAKVLADEVKVKLOKETIDKOCQKIA 708  
QY 196 QLAIVCVSLKKEYENLKEARKASGEVADKLKDLSESSRSKLQTVYSELDOAKLELKSQAK 255  
Db 709 EMVALMEKHKHQYDKIIIEERSELGLYKSKQEOSSLRASLEI---ELSNLKAELLSVKK 765  
QY 256 DLQADKELMSLKKKLMLOETLMLPPVASETVRLVLESPAPVEVNLKLRPSF----- 310  
Db 766 QLEIEREKEKLRKEAR--ENTATLKEKKDKKTQTFLETP--EIIWKLDKSKAVPSQTV 820  
QY 311 -----RDDIDLNATFDVDFP-----PARPSSOHGYEKLCKLEKSH 346  
Db 821 SRNFTSVDHGSKRDKRQYLTWSAKNTLSTPLPKAVTYKTPPKPQLQOR-----ENLN 872  
QY 347 SPIDVPRK 355  
Db 873 IPRESKK 881

RESULT 14  
US-09-310-187A-1  
; Sequence 1, Application US/09310187A  
; Patent No. 6358751  
; GENERAL INFORMATION:  
; APPLICANT: Benichou, Gilles  
; APPLICANT: Fedoseyeva, Eugenia  
; TITLE OF INVENTION: Involvement of Autoantigens in Cardiac  
; TITLE OF INVENTION: Graft Rejection  
; FILE REFERENCE: UCSF-090  
; CURRENT APPLICATION NUMBER: US/09/310,187A  
; CURRENT FILING DATE: 1999-05-12  
; NUMBER OF SEQ ID NOS: 3  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 1  
; LENGTH: 1939  
; TYPE: PRF  
; ORGANISM: Homo sapiens  
US-09-310-187A-1

Query Match 6.5%; Score 156; DB 4; Length 1939;  
Best Local Similarity 22.1%; Pred. No. 0.00015;  
Matches 75; Conservative 59; Mismatches 127; Indels 78; Gaps 10;

QY 79 LKNELDNVRALQSOKDEKRDQOVIIDLRDTLEERNATVVSLOQALGKAEMLCSTLKQ 138  
Db 1289 LAROLEKEKALISOLTRGKLSYTOQMEDLKROLKEEGKAKNALAHALQSAHDDLLR-- 1346  
QY 139 MKYLEQOODETK-----QAQEFAGRLRSKMT-----MQIETLLQSOQLPEV 180  
Db 1347 ----EYEEETLEAKELQRYLSKANSEVAQWRKYETDAIQRTTELEAKKKLAQRQDAE 1402  
QY 181 EEMIRDMGVQSAVEQALAVCVSLK-----EYENLK---EARKASGEVADKLKRDLE 230  
Db 1403 EE-----AVEAVNACSSLEKTKHRLQNETEDLMADVERSNAAAALDKKQNF 1452  
QY 231 SSRSKLQTVYSELDOAKLELKSQAKDLQASADKEIMSLKKL-----TMLQETLMLPV 282  
Db 1453 KILAEWKQKYE---SSELESSOKEARSLSTELFKLKNAYEESLHLETFKRNKNKLOE 1509  
QY 283 VASETVDRVLVLESPAPVEVNLKLRPSFRDDIDLNATFDVDFPAPRPSOHGYEKLCL 342  
Db 1510 EISDLTEQLG-EGGKNVHELEKVRKQLEVEKLEQS-----AL 1546  
QY 343 KSHSPIDVPRKTKCKPKRESQSLGQSCAGPDEEL 381  
Db 1547 EBAEASLEHEBGKILRAQLEFNQIKAEIERKLAKEDEEM 1585

RESULT 15  
US-08-938-105-3  
; Sequence 3, Application US/08938105  
; Patent No. 6353151  
; GENERAL INFORMATION:  
; APPLICANT: Leinwand, Leslie A.  
; APPLICANT: Vikstrom, Karen L.  
; TITLE OF INVENTION: TRANSGENIC MODEL FOR HEART FAILURE  
; NUMBER OF SEQUENCES: 3  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Sheridan Ross P.C.  
; STREET: 1700 Lincoln St., Suite 3500  
; CITY: Denver  
; STATE: CO  
; COUNTRY: U.S.A.  
; ZIP: 80203  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.30  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/938,105  
; FILING DATE:  
; CLASSIFICATION:  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Crook, Marnell M.  
; REGISTRATION NUMBER: 31,071  
; REFERENCE/DOCKET NUMBER: 3595-4  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (303) 863-0223  
; TELEFAX: (303) 863-0223  
; INFORMATION FOR SEQ ID NO: 3:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 1886 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
US-08-938-105-3

Query Match 6.5%; Score 154; DB 4; Length 1886;  
Best Local Similarity 20.6%; Pred. No. 0.00021;  
Matches 82; Conservative 78; Mismatches 148; Indels 90; Gaps 13;

QY 28 TFIHLQCLQSFETAPRTQCRIOYGKRTI---INKLFFDLAQEEENVID-----REFLK 80  
Db 1181 TSHMEQITKA-----KANLEKVSRTLEDQANEYRVKLEEAQORSINDFTTORAKQ 1230  
QY 81 NELDNVRALQSOKD-----KERRDSQOVIIDLRDTLEE---RNAATVVSLOQALGKAE 129  
Db 1231 TENGELAROLEKEKALIMQITRKLSTYTOQMEDLKROLKEEGKAKNALAHALQSAHDD 1290  
QY 130 MLCSTLKKQMKYLEQOODETKQAQEFAGRLRSKMT-----MQIETLLQSOQLPEV 181  
Db 1291 LIREQYEEMEAFAELQRYLSKANSEVAQWRKYETDAIQRTTELEAKKKLAQRQDAE 1350  
QY 182 EEMIRDMGVQSAVEQALAVCVSLK-----EYENLK---EARKASGEVADKLKRDLE 231  
Db 1351 E-----AVEAVNACSSLEKTKHRLQNETEDLMADVERSNAAAALDKKQNF 1400  
QY 232 SRSLQTVYSELDOAKLELKSQAKDLQASADKEIMSLKKL-----TMLQETLMLPV 283  
Db 1401 ILAEWKQKYE---SSELESSOKEARSLSTELFKLKNAYEESLHLETFKRNKNKLOE 1457  
QY 284 ASETVDRVLVLESPAPVEVNLKLRPSFRDDIDLNATFDVDFPAPRPSOHGYEKLCL 343  
Db 1458 ISDLTEQLG-EGGKNVHELEKVRKQLEVEKLEQS-----ALE 1494  
QY 344 KSHSPIDVPRKTKCKPKRESQSLGQSCAGPDEEL 381  
Db 1495 EBAEASLEHEBGKILRAQLEFNQIKAEIERKLAKEDEEM 1532





QY 302 NLRKRPFRDDIDLNA-TFVDVTPPARPSSOHGYEKLCKLEKSHSPIDVPPKICKP 360  
Db 1693 -LKSRFPQARQDLSIDSLDSCCECTPLS-----ITSKL---P 1728  
QY 361 RKESQSLGQSGCGEDELVGAFPIFVNRNAILGQKQKRRPSESSC-----SKDV 412  
Db 1729 RTQPD---GTSVGPGE-----ASPI-----SQRLPPKVESLESFYTPPIPARSQAP 1771  
QY 413 VRTGFDGLG-----GRTKFIQPTDVMIRPLPVK 441  
Db 1772 LESSLDSLGDFVLDGKRTSARRRTQIINITMTKKLDVE 1812

RESULT 18  
US-08-467-781-4  
; Sequence 4, Application US/08467781  
; Patent No. 5780596  
; GENERAL INFORMATION:  
; APPLICANT: TOLKATLY, GARY  
; APPLICANT: LIDGARD, GRAHAM P  
; TITLE OF INVENTION: NOVEL MALIGNANT CELL TYPE MARKERS OF THE  
; TITLE OF INVENTION: INTERIOR NUCLEAR MATRIX  
; NUMBER OF SEQUENCES: 6  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: TESTA, HURWITZ & THIBEAULT  
; STREET: 125 HIGH STREET  
; CITY: BOSTON  
; STATE: MA  
; COUNTRY: USA  
; ZIP: 02110  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/467,781  
; FILING DATE: 06-JUN-1995  
; CLASSIFICATION: 424  
; ATTORNEY/AGENT INFORMATION:  
; NAME: PITCHER ESQ, EDMUND R  
; REGISTRATION NUMBER: 27,829  
; REFERENCE/DOCKET NUMBER: MTP-013  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (617) 248-7000  
; TELEFAX: (617) 248-7100  
; INFORMATION FOR SEQ ID NO: 4:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 2101 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
US-08-467-781-4

Query Match 6.4%; Score 152; DB 1; Length 2101;  
Best Local Similarity 21.0%; Pred. No. 0.00037;  
Matches 97; Conservative 72; Mismatches 160; Indels 132; Gaps 19;

QY 44 RTCPQCRIO-----VGKRTIINKLFFDLAOGEEENV-----LDREFLKNELDNVRQALSOK- 93  
Db 1421 RTAQOQLRAEKASYAEOLSMKKAHGLAENRGLGERANIGRQFLVELDQAREKRYQEL 1480  
QY 94 -----DKERDSOV-----IIDTLRD-----TLEERNATVVSLSQALGRAEML 131  
Db 1481 AAVRADAETLAELVQRAOSTARELEVMYAKYEGSAKVLEBEROFQEROKLTAQVEEL 1540  
QY 132 CSTL-----KROMKYLEOQODETKOQOEAGRLRSKMTMEQIEILLQOLPEVEEMI 184  
Db 1541 SKRLADSDQASKYQOQKLRKAVQAGSGSQEQAQRFQALNE-----LQAQLSQKQ-- 1591  
QY 185 RDMGVGQSAVEQIALVYCVSLKKEYENLKEARKASGE---VADKIRKDLFFSRSLQTVYS 241

Db 1592 -----AAEHYKLQNEKAKTHYDANKQOQNOELQELRSLEQLOKENKELRAEERLGH 1643  
QY 242 ELDOAKLELSAOKDLOSADKEIMSLKKKLTMLQETLNLPPVASETVDRVLVLESPAPVEV 301  
Db 1644 ELQOAGLKTKEAEQTCNHLTAQVRSLEAQVAHADQOL-----RDGKQVATDA----- 1692  
QY 302 NLRKRPFRDDIDLNA-TFVDVTPPARPSSOHGYEKLCKLEKSHSPIDVPPKICKP 360  
Db 1693 -LKSRFPQARQDLSIDSLDSCCECTPLS-----ITSKL---P 1728  
QY 361 RKESQSLGQSGCGEDELVGAFPIFVNRNAILGQKQKRRPSESSC-----SKDV 412  
Db 1729 RTQPD---GTSVGPGE-----ASPI-----SQRLPPKVESLESFYTPPIPARSQAP 1771  
QY 413 VRTGFDGLG-----GRTKFIQPTDVMIRPLPVK 441  
Db 1772 LESSLDSLGDFVLDGKRTSARRRTQIINITMTKKLDVE 1812

RESULT 19  
US-08-195-487-4  
; Sequence 4, Application US/08195487  
; Patent No. 5783403  
; GENERAL INFORMATION:  
; APPLICANT: TOLKATLY, GARY  
; APPLICANT: LIDGARD, GRAHAM P  
; TITLE OF INVENTION: NOVEL MALIGNANT CELL TYPE MARKERS OF THE  
; TITLE OF INVENTION: INTERIOR NUCLEAR MATRIX  
; NUMBER OF SEQUENCES: 6  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: TESTA HURWITZ & THIBEAULT  
; STREET: 53 STATE STREET  
; CITY: BOSTON  
; STATE: MA  
; COUNTRY: USA  
; ZIP: 02109  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; COMPUTER: IBM PC compatible  
; OPERATING SYSTEM: PC-DOS/MS-DOS  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/195,487  
; FILING DATE:  
; CLASSIFICATION: 435  
; PRIOR APPLICATION DATA:  
; APPLICATION NUMBER: US/07/901,701  
; FILING DATE:  
; ATTORNEY/AGENT INFORMATION:  
; NAME: PITCHER ESQ, EDMUND R  
; REGISTRATION NUMBER: 27,829  
; REFERENCE/DOCKET NUMBER: MTP-013  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 617/248-7000  
; TELEFAX: 617/248-7100  
; INFORMATION FOR SEQ ID NO: 4:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 2101 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
US-08-195-487-4

Query Match 6.4%; Score 152; DB 1; Length 2101;  
Best Local Similarity 21.0%; Pred. No. 0.00037;  
Matches 97; Conservative 72; Mismatches 160; Indels 132; Gaps 19;

QY 44 RTCPQCRIO-----VGKRTIINKLFFDLAOGEEENV-----LDREFLKNELDNVRQALSOK- 93  
Db 1421 RTAQOQLRAEKASYAEOLSMKKAHGLAENRGLGERANIGRQFLVELDQAREKRYQEL 1480  
QY 94 -----DKERDSOV-----IIDTLRD-----TLEERNATVVSLSQALGRAEML 131

```

Db 1481 AAVRADAEFLRAEVOAREASTARELEVMYAKYEGAKYKVLLEERQROEEROKLTAQVEEL 1540
QY 132 CSTL-----KKOMKYLEOQODETKOAOEAGRLSKKMTMEQIELLOSQLEPEVEMT 184
Db 1541 SKKLADSDOASKVQOQOKLNAVQAGEESQOEAORFOAQLNE-----LOAQLSQKEQ-- 1591
QY 185 RDMGVGQSAVEQOLAVYCVSLKKEYENLKEARKASGE---VADKLARKDLSSRSKLTQTVS 241
Db 1592 -----AAEHKKLOMEKAKTHYDAKKQONQELQOLNSLEBOLQENKELRAEAERLGH 1643
QY 242 ELDOAKLELKSADKLOSDADKEIMSLKKLTMLQETLNPVVASFTVDRLVLESPPAVEV 301
Db 1644 ELQOAGLKTKEAQOTGRHLTAQVRSLEAQVAHADQOL-----RDLGKFQVATDA---- 1692
QY 302 NKLRRPSFRDDIDLNA-TFDVDTPPARPSSQGHGYEKLCLKESHSPIDVPPKTKCKGP 360
Db 1693 -LKSREPOAKPOLDLSDLSCEEGTPLS-----ITSKL---P 1728
QY 361 RKESQSLSGQSCAGPEDELVGAPLIFVNNALILGOKOPRRPSSSC-----SKDY 412
Db 1729 RTQPD-----GTSVGPB-----ASPL-----SORLPKVESLSLYFTPIPARSQAP 1771
QY 413 VRTGFDGLG-----GRTKFIOPDVTVMIRPLPYK 441
Db 1772 LESSDLSLGDVFLDSGKRTSARRRTQIINIMTKKLQVE 1812

```

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RESULT 20
US-08-483-924-4
; Sequence 4, Application US/08483924
; Patent No. 5882876
; GENERAL INFORMATION:
; APPLICANT: TOKRATLY, GARY
; APPLICANT: LIDGARD, GRAHAM P
; TITLE OF INVENTION: NOVEL MALIGNANT CELL TYPE MARKERS OF THE
; TITLE OF INVENTION: INTERIOR NUCLEAR MATRIX
; NUMBER OF SEQUENCES: 6
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: TESTA, HURWITZ & THIBEAULT
; STREET: 125 HIGH STREET
; CITY: BOSTON
; STATE: MA
; COUNTRY: USA
; ZIP: 02110
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/483,924
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: PITCHER ESQ, EDMUND R
; REGISTRATION NUMBER: 27,829
; REFERENCE/DOCKET NUMBER: MTP-013
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (617) 248-7000
; TELEFAX: (617) 248-7100
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2101 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-483-924-4

```

Query Match 6.4%; Score 152; DB 2; Length 2101;  
 Best Local Similarity 21.0%; Pred. No. 0.00037;  
 Matches 97; Conservative 72; Mismatches 160; Indels 132; Gaps 19;

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QY 44 RTCPQCRIO---VGKRTIINKLFPDLAQEEENV-----LDREFLKNELDNVRALQSLQK- 93
Db 1421 RTAQQLRAEKASYAEOLSLMKAHGLLAEBNRGLGRANLGRQFLVEYLDQAREKTYQEL 1480
QY 94 -----DKERDSQV---IIDTLRD-----TLEENNAIVVSLQALGKRAEML 131
Db 1481 AAVRADAEFLRAEVOAREASTARELEVMYAKYEGAKYKVLLEERQROEEROKLTAQVEEL 1540
QY 132 CSTL-----KKOMKYLEOQODETKOAOEAGRLSKKMTMEQIELLOSQLEPEVEMT 184
Db 1541 SKKLADSDOASKVQOQOKLNAVQAGEESQOEAORFOAQLNE-----LOAQLSQKEQ-- 1591
QY 185 RDMGVGQSAVEQOLAVYCVSLKKEYENLKEARKASGE---VADKLARKDLSSRSKLTQTVS 241
Db 1592 -----AAEHKKLOMEKAKTHYDAKKQONQELQOLNSLEBOLQENKELRAEAERLGH 1643
QY 242 ELDOAKLELKSADKLOSDADKEIMSLKKLTMLQETLNPVVASFTVDRLVLESPPAVEV 301
Db 1644 ELQOAGLKTKEAQOTGRHLTAQVRSLEAQVAHADQOL-----RDLGKFQVATDA---- 1692
QY 302 NKLRRPSFRDDIDLNA-TFDVDTPPARPSSQGHGYEKLCLKESHSPIDVPPKTKCKGP 360
Db 1693 -LKSREPOAKPOLDLSDLSCEEGTPLS-----ITSKL---P 1728
QY 361 RKESQSLSGQSCAGPEDELVGAPLIFVNNALILGOKOPRRPSSSC-----SKDY 412
Db 1729 RTQPD-----GTSVGPB-----ASPL-----SORLPKVESLSLYFTPIPARSQAP 1771
QY 413 VRTGFDGLG-----GRTKFIOPDVTVMIRPLPYK 441
Db 1772 LESSDLSLGDVFLDSGKRTSARRRTQIINIMTKKLQVE 1812

```

```

RESULT 21
US-09-452-294-1
; Sequence 1, Application US/09452294
; Patent No. 6287790
; GENERAL INFORMATION:
; APPLICANT: Delievre, Sophie
; APPLICANT: Bissell, Mina
; TITLE OF INVENTION: UTILIZATION OF NUCLEAR STRUCTURAL PROTEINS FOR TARGETED
; TITLE OF INVENTION: THERAPY AND DETECTION OF PROLIFERATIVE AND
; FILE REFERENCE: IB-1454- Sequence Submittal
; PATENT REFERENCE: IB-1454- Sequence Submittal
; CURRENT APPLICATION NUMBER: US/09/452,294
; CURRENT FILING DATE: 1999-11-30
; PRIOR APPLICATION NUMBER: 60/110,420
; PRIOR FILING DATE: 1998-11-30
; NUMBER OF SEQ ID NOS: 1
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 2101
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-452-294-1

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Query Match 6.4%; Score 152; DB 4; Length 2101;  
 Best Local Similarity 21.0%; Pred. No. 0.00037;  
 Matches 97; Conservative 72; Mismatches 160; Indels 132; Gaps 19;

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QY 44 RTCPQCRIO---VGKRTIINKLFPDLAQEEENV-----LDREFLKNELDNVRALQSLQK- 93
Db 1421 RTAQQLRAEKASYAEOLSLMKAHGLLAEBNRGLGRANLGRQFLVEYLDQAREKTYQEL 1480
QY 94 -----DKERDSQV---IIDTLRD-----TLEENNAIVVSLQALGKRAEML 131
Db 1481 AAVRADAEFLRAEVOAREASTARELEVMYAKYEGAKYKVLLEERQROEEROKLTAQVEEL 1540
QY 132 CSTL-----KKOMKYLEOQODETKOAOEAGRLSKKMTMEQIELLOSQLEPEVEMT 184

```

Db 1541 SKRLADSDQASKVQOQKLVKAVQAGSGEOEAFQFQALNE-----LQAOISQKEQ-- 1591  
QY 185 RDMGVQGSAAVEQALAVYCVSLKKEYENLKEARKASGE---VADKLKRDLPSSRSKLTQTVS 241  
Db 1592 -----AAEHKKLQMEKAKTHTYDAKKQONQELQEQRLSLEQLOQENKELRAEAERLGH 1643  
QY 242 ELDOAKLELKSAAKDLQASADKEIMSLKKLTMLQETLNLPPVASETVDRVLVLESPARVEV 301  
Db 1644 ELQOAGLKTKEAEQOTCHHLTAQVRSLEAQVAHADQOL-----RDLGKFOVATDA----- 1692  
QY 302 NLKLRSPFDDIDLNA-TDVDTPPARPSSSHGYEKLCKLEKSHSPIQDVKKICKGP 360  
Db 1693 -LKSREPQAKPQDLSDLSIDLSCBECTPLS-----RDLGKFOVATDA----- 1728  
QY 361 RKESQSLGQSCAGPEDEELVGAFPIFYRNAILGOKOPKRPRESSC-----SKDV 412  
Db 1729 RTQPD-----GTSVGP-----ASPI-----SQRLLPKVESLESLEYFTPIPARSQAP 1771  
QY 413 VRTGFDGLG-----GRTKFIQPTDVTVMIRPLPVK 441  
Db 1772 LESSLDLGDVFLDGSGRKTRRSARRRTQIINITMTKKLDVE 1812

RESULT 22  
PCT-US93-06160-4  
; Sequence 4, Application PC/TUS9306160  
; GENERAL INFORMATION:

; APPLICANT:  
; TITLE OF INVENTION: NOVEL MALIGNANT CELL TYPE MARKERS OF THE  
; TITLE OF INVENTION: INTERIOR NUCLEAR MATRIX  
; NUMBER OF SEQUENCES: 6  
; CORRESPONDENCE ADDRESSES:  
; ADDRESSEE: TESTA HURWITZ & THIBEAULT  
; STREET: 53 STATE STREET  
; CITY: BOSTON  
; STATE: MA  
; COUNTRY: USA  
; ZIP: 02109  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; OPERATING SYSTEM: IBM PC compatible  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: PCT/US93/06160  
; FILING DATE: 19930621  
; CLASSIFICATION:  
; ATTORNEY/AGENT INFORMATION:  
; NAME: PITCHER ESQ, EDMUND R  
; REGISTRATION NUMBER: 27,829  
; REFERENCE/DOCKET NUMBER: MTP-013  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: 617/248-7000  
; TELEFAX: 617/248-7100  
; INFORMATION FOR SEQ ID NO: 4:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 2101 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
PCT-US93-06160-4

Query Match 6.4%; Score 152; DB 5; Length 2101;  
Best Local Similarity 21.0%; Pred. No. 0.00037;  
Matches 97; Conservative 72; Mismatches 132; Gaps 19;

QY 44 RTCPQCRIQ---VGRRTIINKLFFDLAEEENV-----LDREFLKNELDNVPRAQLSQK- 93  
Db 1421 RTMQOLRAEKASYAEQLSLMKRAHGLLAEEENRGLGRANLGRFLVELDQAREKXYQEL 1480  
QY 94 -----DKERKDSQV---IIDTLRD-----TLEERNATVVSILQALGKAEML 131

Db 1481 AAVRADAEETRLAEVREAOASTARELEVMPTAKYEGAKVYLEERQRFQOEROKLTLAQVEEL 1540  
QY 132 CSTL-----KKOMKYLBDQOQDETQAOEAGRLSKSKMTNQIELLOSLQPEVEEMI 184  
Db 1541 SKRLADSDQASKVQOQKLVKAVQAGSGEOEAFQFQALNE-----LQAOISQKEQ-- 1591  
QY 185 RDMGVQGSAAVEQALAVYCVSLKKEYENLKEARKASGE---VADKLKRDLPSSRSKLTQTVS 241  
Db 1592 -----AAEHKKLQMEKAKTHTYDAKKQONQELQEQRLSLEQLOQENKELRAEAERLGH 1643  
QY 242 ELDOAKLELKSAAKDLQASADKEIMSLKKLTMLQETLNLPPVASETVDRVLVLESPARVEV 301  
Db 1644 ELQOAGLKTKEAEQOTCHHLTAQVRSLEAQVAHADQOL-----RDLGKFOVATDA----- 1692  
QY 302 NLKLRSPFDDIDLNA-TDVDTPPARPSSSHGYEKLCKLEKSHSPIQDVKKICKGP 360  
Db 1693 -LKSREPQAKPQDLSDLSIDLSCBECTPLS-----RDLGKFOVATDA----- 1728  
QY 361 RKESQSLGQSCAGPEDEELVGAFPIFYRNAILGOKOPKRPRESSC-----SKDV 412  
Db 1729 RTQPD-----GTSVGP-----ASPI-----SQRLLPKVESLESLEYFTPIPARSQAP 1771  
QY 413 VRTGFDGLG-----GRTKFIQPTDVTVMIRPLPVK 441  
Db 1772 LESSLDLGDVFLDGSGRKTRRSARRRTQIINITMTKKLDVE 1812

RESULT 23  
US-08-095-737-2  
; Sequence 2, Application US/08095737  
; Patent No. 5487979  
; GENERAL INFORMATION:

; APPLICANT: DiFiore, Pier P  
; TITLE OF INVENTION: A Substrate for the Epidermal Growth  
; TITLE OF INVENTION: Factor Receptor Kinase  
; NUMBER OF SEQUENCES: 4  
; CORRESPONDENCE ADDRESSES:  
; ADDRESSEE: Knobbles, Martens, Olson & Bear  
; STREET: 620 Newport Center Drive, Sixteenth Floor  
; CITY: Newport Beach  
; STATE: California  
; COUNTRY: United States of America  
; ZIP: 92660  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Floppy disk  
; OPERATING SYSTEM: IBM PC compatible  
; SOFTWARE: PatentIn Release #1.0, Version #1.25  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/095,737  
; FILING DATE: 19930722  
; CLASSIFICATION: 530  
; ATTORNEY/AGENT INFORMATION:  
; NAME: Israelisen, Ned A  
; REGISTRATION NUMBER: 29,655  
; REFERENCE/DOCKET NUMBER: NIH060.001A  
; TELECOMMUNICATION INFORMATION:  
; TELEPHONE: (619) 235-8550  
; TELEFAX: (619) 235-0176  
; INFORMATION FOR SEQ ID NO: 2:  
; SEQUENCE CHARACTERISTICS:  
; LENGTH: 896 amino acids  
; TYPE: amino acid  
; TOPOLOGY: linear  
; MOLECULE TYPE: protein  
US-08-095-737-2

Query Match 6.2%; Score 149; DB 1; Length 896;  
Best Local Similarity 22.8%; Pred. No. 0.0002;  
Matches 62; Conservative 46; Mismatches 110; Indels 54; Gaps 8;

QY	17	SRDVAALHCIGTFIL- : : : : : : : : : : : : : : :	OCLOSFE-----	TAPSTQCQCRIQYKGRITINIKLFEFL	AOE	69
Db	278	SKDOFAL-----	AFHLISOKLIKIGDIPHVLTPEKIMPPSDRASLOKNITIGSSPAVDSAIK			333
QY	70	EENVILDRFF-LKNELDNVRAQLSQDKDERDSOVIIIDTLRDTL	EEHNATVSLQALGK			127
Db	334	ELDTLNNEIYDLOREKNNVEODLKEKEDTJKQRTSEVODLODEVQRENTNLOKLOAKOO				393
QY	128	AEMCTSLKKOMKYLEOQOETKQAGEARLSKMTQOIELLOQSLPEVPEEMIRDM				187
Db	394	VOELLDLDEKQAOLEQLEKVRKKCAEAPOLISLSLA-----	ELTJSE-----			437
QY	188	GVGSAVEQJLAUVYCVSLKEKEYENLKEARKASGEVADTLRKDLDFSSRSKLOTQVYSELDOAK				247
Db	438	-----SQISTVEEELAKKRREELSRLOQETALEE-----	SVESGK			472
QY	248	LELSAKQDIQASAKDEIMSLKKRTITMQOETFLN				279
Db	473	AOLEPDLQOHDQSOEILSSMOQKLEMKKMDLEN				504

RESULT 24  
US-08-480-145-2

MOLECULE TYPE: protein  
US-08-480-145-2

OY	70	EEVYLDRE---LKNELNVAQLOSKOKERKRDSDVITLDTLDTLEENNAVSVLSLOALG	127
Db	334	ELDTPLNNEIYDLQEKKNNEVDOLKEKEDDTIKQRTSEVQDLODEYQRENTMQLQLOAKQO	333
OY	128	AEMLCSTLKKOMKYLLEQOODETRKQOEAGRLRSKMTMEQIIEILLQOLPEVEEMIRDM	187
Db	394	VQELDELDEBQKAQLEBQLKEVRKKCAEBMQLISLKA---ELTSGE	437
OY	188	GVGSAVEQLAVYCVSLKKELENLKEKERRKASGEVADKLKRDLFSSRSKLOTVYSELDQAK	247
Db	438	-----SQISTVEEELAKAREELSRQOETAELEE-----SVESGK	472
OY	248	LELSAQKDLOSADEKELSLTKKLTMLQOETLN	279
Db	473	AOLEPQOHLQDQSQOETISSMOKMKEMKQDEN	504

RESULT 25  
US-08-477-389-2  
; sequence 2, Application US/08477389

MOLECULE TYPE: protein  
US-08-477-389-2

```
Qy 17 SRDVAIHCGHTFHL--QCLDSE----TAPSTCPQCRQVKGRTIINKLFEDL-AQE 69
      | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 278 SKDQFAL----AFHLISQKLKIGIDPPhVLTPEMIIPSDRASLQNNIGSSPVDAFSAIK 333
```

Db 334 ELDTLNNELVLDQREKNVNEODLKEKEDTJKQRTSEVODLQDEVQRENTNLQKLOAQKQ 393  
QY 128 AEMCLSTLKKOMKYLBOQOQDTKQAOEAGRLRSKMTMEIILLQSOLPEVEEMIRDM 187  
Db 394 VOELLDELDEKQALQEROLKEVRKKCAEADLISSLSKA-----ELTSE----- 437  
QY 188 GVGSAVEQJLAVCVSLKREYENLEKARKASGEVADLRKDLFSSRSKLTQVYSELDOAK 247  
Db 438 -----SQISTVEELAKAREELSRLQOETAELEE-----SVESGK 472  
QY 248 LEKSAQKDLQASADKETMSLKKLTLMQETLN 279  
Db 473 AQLEPLQOHLDQSOEISMQMKLMEMKLDLEN 504

RESULT 26  
PCT-US93-03077-1  
Sequence 1, Application PC/TUS9303077  
GENERAL INFORMATION:  
APPLICANT: Board of Regents, The University of Texas System  
APPLICANT: Gaynor, Richard B.  
APPLICANT: Wu, Foon Kin  
TITLE OF INVENTION: PROTEIN CELLULAR FACTOR USEFUL FOR  
TITLE OF INVENTION: REGULATING GENE EXPRESSION  
NUMBER OF SEQUENCES: 7  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Arnold White & Durkee  
STREET: P.O. Box 4433  
CITY: Houston  
STATE: Texas  
COUNTRY: USA  
ZIP: 77210

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: PCT/US93/03077  
FILING DATE: 19930331  
CLASSIFICATION:  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US/07/862,025  
FILING DATE: April 2, 1992  
ATTORNEY/AGENT INFORMATION:  
NAME: Kammerer, Patricia A.  
REGISTRATION NUMBER: 29,775  
REFERENCE/DOCKET NUMBER: UFD270PCT  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 713-787-1540  
TELEFAX: 713-749-2679  
TELEX:  
INFORMATION FOR SEQ. ID NO: 1:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 1093 amino acids  
TYPE: AMINO ACID  
STRANDEDNESS: unknown  
TOPOLOGY: unknown  
MOLECULE TYPE: protein  
PCT-US93-03077-1

Query Match 6.2%; Score 149; DB 5; Length 1093;  
Best local Similarity 21.3%; Pred. No. 0.00076;  
Matches 99; Conservative 88; Mismatches 155; Indels 122; Gaps 21;

QY 59 INKLFDFLAQEEEN--VLD--REFLNEDLVNRAQTSQKDKERD--SQYITIDLRPTLE 112  
Db 593 LNKVKELEELLOHKLQVLDGKREVEQKHENIKLINSWVERQKDLGRLOVD--HDELE 650  
QY 113 ERNATVVSLOALGKA-----EMLCSTLKKOMK-----YLEQOQDET 149  
Db 651 EKNR-----SIGAALDSAVKELTDLHKANAKDSQAQALSLREMKAKBELSALEKQGEA 707

QY 150 KQAGE-----EAGRLRSKMTMEIELL-----LOSOLPEVEEMIRDMGVGSAY 194  
Db 708 KQOQETLAIQVGDURLALQRTQEAARKEDYLRIHEITGLOQDAENRQDSQSVST 767  
QY 195 EQLAVVCSLKKREYENLEKARKASGEVADLRKDLFSSRSKLTQVYSELDOAKLELSAQ 254  
Db 768 TR-----PLLRQENLQATLGISQTSMEKLEKLSRLGESQL-----LAAAV 811  
QY 255 KDLQASAKETMSLKKLTLMQETLNLPVASEYDRLVLESAPAEVNLKLRPSFDDI 314  
Db 812 ERERAATEELLANKIQMSM-----ESONSILROENSFRQALSEKNRLCKLE 860  
QY 315 DLNATFVDVPPAPRSSQGGYKELCLEKSHSPIDVPPKICKGPKKE-----SOLSLG 369  
Db 861 DENNRIOVE-----LEN-----LKDEYVRLTEETREKXTLNSOLEME 898  
QY 370 GQSCAGEPDEELVGAFFIFRNAILGOKRPRSESS---CSKDVYRTGFDGIGRTKF 426  
Db 899 RMKVEGRKK-----ALFTQETI--KEKERKPPSVSSTPTMSRSSISGVDMAGLQTSF 950  
QY 427 IQPTDT--VAIRPLVPVPRKTK-----VKQRYKTVTSLEQAKL 463  
Db 951 LSODESHDSFGPMPISAKMKHLYACKDGSRIKHIEML-QSOL 993

RESULT 27  
US-08-687-080-51  
Sequence 51, Application US/08687080  
Patent No. 5965427  
GENERAL INFORMATION:  
APPLICANT: Gregory Dolganov  
TITLE OF INVENTION: Human RAD50 Gene and Methods of Use Thereof  
NUMBER OF SEQUENCES: 175  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Dehlinger & Associates  
STREET: 350 Cambridge Avenue, Suite 250  
CITY: Palo Alto  
STATE: CA  
COUNTRY: USA  
ZIP: 94306

COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/687,080  
FILING DATE: 17-JUL-1996  
CLASSIFICATION: 435  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US 08/592,126  
FILING DATE: 26-JAN-1996  
ATTORNEY/AGENT INFORMATION:  
NAME: Sholtz, Charles K.  
REGISTRATION NUMBER: 38,615  
REFERENCE/DOCKET NUMBER: 4600-0111.30  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (415) 324-0880  
TELEFAX: (415) 324-0960  
INFORMATION FOR SEQ. ID NO: 51:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 1312 amino acids  
TYPE: amino acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
HYPOTHEICAL: NO  
ANTI-SENSE: NO  
ORIGINAL SOURCE:  
INDIVIDUAL ISOLATE: TRANS. OF RAD50 cDNA (SEQ. 54), NT.  
INDIVIDUAL ISOLATE: 389 TO 4324  
US-08-687-080-51



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RESULT 31
US-08-592-126-148
: Sequence 148, Application US/08592126
: Patent No. 5821091
: GENERAL INFORMATION:
: APPLICANT: Gregory Dolganov
: TITLE OF INVENTION: Transcripts Encoding Immunomodulatory
: TITLE OF INVENTION: Polypeptides
: NUMBER OF SEQUENCES: 151
: CORRESPONDENCE ADDRESS:
: ADDRESSEE: Dehlinger & Associates
: STREET: 350 Cambridge Avenue, Suite 250
: CITY: Palo Alto
: STATE: CA
: COUNTRY: USA
: ZIP: 94306
: COMPUTER READABLE FORM:
: MEDIUM TYPE: Floppy disk
: COMPUTER: IBM PC compatible
: OPERATING SYSTEM: PC-DOS/MS-DOS
: SOFTWARE: PatentIn Release #1.0, Version #1.25
: CURRENT APPLICATION DATA:
: APPLICATION NUMBER: US/08/592,126
: FILING DATE:
: CLASSIFICATION: 435

```



ATTORNEY/AGENT INFORMATION:  
NAME: Sholtz, Charles K.  
REGISTRATION NUMBER: 38,615  
REFERENCE/DOCKET NUMBER: 4600-0111  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (415) 324-0880  
TELEFAX: (415) 324-0960  
INFORMATION FOR SEQ ID NO: 148:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 1312 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
HYPOTHETICAL: NO  
ORIGINAL SOURCE:  
INDIVIDUAL ISOLATE: Rad50.pro-translation of SEQ ID NO:54  
US-08-592-126-148

Query Match 6.2%; Score 147.5; DB 2; Length 1312;  
Best Local Similarity 22.2%; Pred. No. 0.00046;  
Matches 98; Conservative 69; Mismatches 126; Indels 149; Gaps 22;

QY 52 QVCKRTITKLFPLAOF-----ENVLDR-----EFL---KNELDNVRAQ 89  
DB 404 QCEEAKTANQIMNDFAEKETLKOKQIDETDKKTGLRIITELKSEILSKKNELKNVKE 463  
QY 90 LSG-----KDKERKDSQVIIDTLRD-TLEERNAT-----VSLQALGKAEV----- 130  
DB 464 LQLEGSSDRILDELDELKAERELSKAEKNSVETLKMEVLSLQNE--KADIDRFLRL 521  
QY 131 -----LCSTL-----KKQMK-YLE 143  
DB 522 DQEMEDLNHTTTRTQKEMLTQKADKDEQIRIKSHSDELTLGLGFNNKQLEWMLH 561  
QY 144 QQODETKQAEERGLRSKKMTMEQIELLIQSOLPEVEEMIRDMVGQSAVEQLAYCVS 203  
DB 582 SKSEKINQTRDLAKLNEKLASSSEONKNHINELKREEOI-----SSYEDKLFVCGS 635  
QY 204 -----LKKEVE-NLKERKASG-----EVADK-----LRKDLSSRSKIQ 237  
DB 636 QDFESDLRKEETEKSSKORAMLAGATAVYSQFTQLTENOSCCPVCORVQTELEIQ 695  
QY 238 TVSELDQAKLEKSAQKDSADKEIMSLKKLTMLQETLNPVASEFVDRLVESPA 297  
DB 696 EYISDL-QSKLRL--APDKLSTSE--LKKEKRDMLGLVPMQSIIDLEKEKIP- 748  
QY 298 PVEVNLKLRPS-----FRDDILNATFVDVTPPARPSSOHGYEKLCT-----EKSH 346  
DB 749 --ELRNKLQVNRDIOQLKNDIEOEFTLGLTIMPEESA-----KVCULTDVTIMRFQ 799  
QY 347 SPIODVKKICKGPKRESQSL 368  
DB 800 MELKDVERRKTAQQAQKLGIDL 821

## RESULT 32

US-09-085-199B-4  
Sequence 4, Application US/09085199B  
Patent No. 6235879  
GENERAL INFORMATION:  
APPLICANT: Hayden, Michael R.  
APPLICANT: Hackam, Abigail  
APPLICANT: Hug, A.H.M. Mahubul  
APPLICANT: Chopra, Vikramjit Singh  
APPLICANT: Kalchman, Michael  
TITLE OF INVENTION: Apoptosis Modulators That Interact with the  
TITLE OF INVENTION: Huntington's Disease Gene  
NUMBER OF SEQUENCES: 44  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Opedahl & Larson  
STREET: PO Box 5270  
CITY: Frisco

STATE: CO  
COUNTRY: USA  
ZIP: 80443-5270  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 KB storage  
COMPUTER: IBM Compatible  
OPERATING SYSTEM: MS DOS 5.0  
SOFTWARE: Wordperfect  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/085,199B  
FILING DATE:  
CLASSIFICATION:  
ATTORNEY/AGENT INFORMATION:  
NAME: Larson, Marina T.  
REGISTRATION NUMBER: 32038  
REFERENCE/DOCKET NUMBER: UBC-P-013052  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (970) 668-2050  
TELEFAX: (970) 668-2052  
INFORMATION FOR SEQ ID NO: 4:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 914  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
HYPOTHETICAL: no  
ORIGINAL SOURCE:  
ORGANISM: human  
FEATURE: Huntington-interacting protein  
US-09-085-199B-4

Query Match 6.1%; Score 146.5; DB 4; Length 914;  
Best Local Similarity 21.1%; Pred. No. 0.00033;  
Matches 54; Conservative 63; Mismatches 92; Indels 47; Gaps 8;

QY 59 INKLFEDLAQEEENVLDREFLKNELDNVRAQLSQDKERKDSQVIIDTLRDITLEERNATV 118  
DB 245 VNK-----DEKHLIER--LYREISGLKQI---ENMKTESQRVVYLQKGHVSELEADL 293  
QY 119 VSLQALGKAEMLCSTLKKOMKYLEQODETKQAEERGLRSKKMTMEQIELLIQSOLP 178  
DB 294 AEOQHRLQQAADCEFLRLDELDELRRQREDTEKAQRSLSIEIRKAQANEQRYSLKEKYS 353  
QY 179 EY-----EEMIRDMVGQSAVEQLAYCVSLKREYENLKEA-----RKASG--EVD 223  
DB 354 ELVQNHADLKNKAEVTKOVSMARQAVDLEREKKELEDSLERISPOGOKTKTOEQLVLE 413  
QY 224 KLRKDLFSSRSKIQTV-----YSELQAKLEKSA---QKDLQADK 262  
DB 414 SLKQELGTSQRELQVIGSLSTSAQSEANMAAFELEREKRDLSVGAHREBELSALK 473  
QY 263 EIMSLKKLTMLQETL 278  
DB 474 ELQDTQLKLASTEEM 489

## RESULT 33

US-09-085-199B-11  
Sequence 11, Application US/09085199B  
Patent No. 6235879  
GENERAL INFORMATION:  
APPLICANT: Hayden, Michael R.  
APPLICANT: Hackam, Abigail  
APPLICANT: Hug, A.H.M. Mahubul  
APPLICANT: Chopra, Vikramjit Singh  
APPLICANT: Kalchman, Michael  
TITLE OF INVENTION: Apoptosis Modulators That Interact with the  
TITLE OF INVENTION: Huntington's Disease Gene  
NUMBER OF SEQUENCES: 44  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Opedahl & Larson  
STREET: PO Box 5270

CITY: Frisco  
STATE: CO  
COUNTRY: USA  
ZIP: 80443-5270  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Kb storage  
COMPUTER: IBM Compatible  
OPERATING SYSTEM: MS DOS 5.0  
SOFTWARE: WordPerfect  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/09/085,199B  
FILING DATE:  
CLASSIFICATION:  
ATTORNEY/AGENT INFORMATION:  
NAME: Larson, Marina T.  
REGISTRATION NUMBER: 32038  
REFERENCE/DOCKET NUMBER: UBC-P-013052  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (970) 668-2050  
TELEFAX: (970) 668-2052  
INFORMATION FOR SEQ ID NO: 11:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 1068  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
HYPOTHEICAL: no  
ORIGINAL SOURCE:  
ORGANISM: mouse  
FEATURE:  
OTHER INFORMATION: Huntington-interacting protein -mHipla  
US-09-085-199B-11

Query Match 6.1%; Score 146; DB 4; Length 1068;  
Best Local Similarity 21.8%; Pred. No. 0.00046;  
Matches 113; Conservative 84; Mismatches 187; Indels 134; Gaps 23;  
QY 30 HLOCILQSPFAPSRTPCPOCRIOY-----GKRTIINKLFFDL-----AOEENVDRE 77  
DB 298 HIRVYVYIPPEAPPEEPENLIESSAPPGEPYVADLDOTGPPNGSKDDRIQIE 357  
QY 78 FLKNEIDNVAQLSQDKERKDSQVIIDTLRD-----TLEERNATVLSQALGKRAEM 130  
DB 358 NIKREVEYTLRAEL--EKIKMEARQYISQLKGVNGLEAELEBQR--QKRALVDNEQ 411  
QY 131 LCSTLKQMKYLEQOODETQAOEEAGR-----LRSKMK 164  
DB 412 LRHEL-AQLALDLEGARRNGLREERKASATARYSKLEKHSINTHAEILRNAD 470  
QY 165 TMEQIELLQSQLPVEEMTRDMGVGSAVEQLAVYCVSLKKEVENLKEARKASGEVADK 224  
DB 471 TAKQLVYTOOSO--EVAR-----YKQLAFQMEQAKRESKMEQ--SDLEK 516  
QY 225 LRDLFSSRSKL--QTVSELDOAKLELSAOKDLOSADKEIMS--LKKRLTMLQETLN 279  
DB 517 LKRELAARAGELARAQALSRTQSGSEL--SSRLDTLNARKKALSGVVRREARL----- 570  
QY 280 LRPVASTVTRVLVESAPEVNLKLRPSFRDDIDLNATFDVDTPPARPSSSOHGYEK 339  
DB 571 --LAQSLVREKEEALSQOQRSSQKGLRQL-----AEKESQEOGLRQK 615  
QY 340 LCLE-----KSHSPIDGVPKKICKGPKRESOLSLGSGSCAGPEDELGAPPIFV 389  
DB 616 LDEQLAVLRSAAEAAEAILIQDAVSKL-----DDPLHL--RCTSSPDILVSRQAAL- 655  
QY 390 RNAILQOKQPKR--PRSESSCKDVTYRTGDFL-----GGRTKFIQPTDTV----- 433  
DB 666 -DSVSGLEGCHTQYGLASSSEDAALVALTRFSLHADTVINGAATSHLATPDADRLMDT 724  
QY 434 ----MIRPLPVKPKTKYKQVRYKYVPSLFQAKLDITFL 467  
DB 725 CRECGAAELVGLQ-QDQTVLRRAQPSLMRAPLQGITL 761

RESULT 34  
US-08-600-982-24  
Sequence 24, Application US/08600982  
Patent No. 6120991  
GENERAL INFORMATION:  
APPLICANT: Carter, William G.  
APPLICANT: Gil, Susana A.  
APPLICANT: Ryan, Maureen C.  
TITLE OF INVENTION: Epiligrin, an Epithelial Ligand for  
NUMBER OF SEQUENCES: 30  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Christensen, O'Connor, Johnson, and Kindness  
STREET: 1420 Fifth Avenue  
CITY: Seattle  
STATE: WA  
COUNTRY: USA  
ZIP: 98101-8100  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/600,982  
FILING DATE: 02-SEP-1994  
CLASSIFICATION: 435  
ATTORNEY/AGENT INFORMATION:  
NAME: Shelton, Dennis K.  
REGISTRATION NUMBER: 26,997  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (206) 682-8100  
TELEFAX: (206) 224-0779  
INFORMATION FOR SEQ ID NO: 24:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 1713 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
DESCRIPTION: E170 protein as translated from sequence  
DESCRIPTION: of FIGURES 15A-15F, and as shown also in FIGURES  
DESCRIPTION: 19A-19R  
US-08-600-982-24  
Query Match 6.1%; Score 145.5; DB 3; Length 1713;  
Best Local Similarity 20.7%; Pred. No. 0.00099;  
Matches 85; Conservative 55; Mismatches 116; Indels 155; Gaps 17;  
QY 34 LIQSFETAPSRTPCPOCR-----IQVGRKTIINKLFFDLQO-----E 69  
DB 278 LNOEFETLQEKAAVNSRKQATLNNVNRATQSAKELDVKIKVIRNVHILLQISGTGE 337  
QY 70 EENVVLDREF-----LKNELDNVRAQLSQDKERKDSQYII----- 104  
DB 338 GNNVPSGDFSREMAEQRMRELNR--NFGKLRLEADKRESQULLRIRTWOKTHG 395  
QY 105 ----DTLRDLTLEERNATV--VSLQALGKA-----EMLCSTLKQMKYLEQ 144  
DB 396 ENGNLANISIRDSINEYEAKLSIDLRARLQEAQAQANGLNQENEFALGAIORVKEIMS 455  
QY 145 -QODEK-----QAOEAGRLRSKMKTKTEQJLELLQSLPVEEMTRDM--GVQO 191  
DB 456 LOSDFPKYLTADSSLLQTNIALQIMEKSOKEVEKLAASLNEARQSLSKVRELSRSAGK 515  
QY 192 -SAVEQLAVYCVSLK-----EYENLKARKASGEVADK 224  
DB 516 TSLVEAEKHAASLOELAOLEIRKRNAGSDELYRCVADAATVETILNAIAAEBAANR 575  
QY 225 LRDLFSSRSKLQTYVSE-----LDQAKLELSAOKDLOSADKEIMSLK 268

Db 576 AAS---ASESALQTVIKEDLPKRAKTLSSNSDKLNEAKMTOKLKQEVSPA----- 624

Qy 269 KKLTMLOETLNPVASETVDRVLVLESPAPVEYNLKLRRPSF-----RDDID 315

Db 625 --LNNIQOTLNTIVTVQKEVID-----TNLTTLRLDGLHGIGRGDID 662

RESULT 35

PCT-US94-10261A-24

Sequence 24, Application PC/TUS9410261A

GENERAL INFORMATION:

APPLICANT: Carter, William G.

APPLICANT: Gil, Susanna A.

APPLICANT: Ryan, Maureen C.

TITLE OF INVENTION: Epiligrin, an Epithelial Ligand for

TITLE OF INVENTION: Integrins

NUMBER OF SEQUENCES: 30

CORRESPONDENCE ADDRESS:

ADDRESSEE: Christensen, O'Connor, Johnson, and Kindness

STREET: 1420 Fifth Avenue

CITY: Seattle

STATE: WA

COUNTRY: USA

ZIP: 98101-8100

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: PCT/US94/10261A

FILING DATE: 02-SEP-1994

CLASSIFICATION:

ATTORNEY/AGENT INFORMATION:

NAME: Shelton, Dennis K.

REGISTRATION NUMBER: 26,997

TELECOMMUNICATION INFORMATION:

TELEPHONE: (206) 682-8100

TELEFAX: (206) 224-0779

INFORMATION FOR SEQ. ID NO: 24:

SEQUENCE CHARACTERISTICS:

LENGTH: 1713 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

DESCRIPTION: E170 protein as translated from sequence of

FIGURES 15A-15F, and as shown also in FIGURES 19A-19R

PCT-US94-10261A-24

Query Match 6.1%; Score 145.5; DB 5; Length 1713;

Best Local Similarity 20.7%; Pred. No. 0.00099;

Matches 85; Conservative 55; Mismatches 116; Indels 155; Gaps 17;

Qy 34 LQSFFETAPSRTPQPCR-----IOVGRRTIINKLFFDLAQ-----E 69

Db 278 LNFQEFLLQKKAQYNSKQATLNNVNRATQSAKELDVKIKNVIRNVHILLKQISGTGDE 337

Qy 70 EENVLDREF-----LKNELDNVRAQLSQDKKERSQVYI----- 104

Db 338 GNNVPSDGFREWNAEQARMKRELNR--NFGKHLREAEADKRRESQLLNIRITVQKTHQG 395

Qy 105 -----DTLDDTLERRNAVY-----VSLQALGRA-----EMLCSTLKKQMKYLEQ 144

Db 396 ENNGCLASIRDSLNEYEAKLSDLRARLQEAQAQKQANGLENRERALGAIQROVKEINS 455

Qy 145 -QODETK-----QAQDEAGRLRSKMKTYEQIELLQSOLPEVEEMIRDW--GVGQ 191

Db 456 LQSFYTYLTATDSSLQTNIALQLMKRSQKEYEKLASLNEARQELSDVKYRELSTRAGK 515

Qy 192 -SAVEQLAVYCVSLK-----EYENLKEARKASGEVADK 224

Db 516 TSLVEAEKHAHSIQELAKOLEEIKRNASGDELVCADVADATAYENILNKAEDANR 575

Qy 225 LRKDLFSSRSKLOTYXSE-----LDOAKLELKSQKDLQSADEIMSLK 268

Db 576 AAS---ASESALQTVIKEDLPKRAKTLSSNSDKLNEAKMTOKLKQEVSPA----- 624

Qy 269 KKLTMLOETLNPVASETVDRVLVLESPAPVEYNLKLRRPSF-----RDDID 315

Db 625 --LNNIQOTLNTIVTVQKEVID-----TNLTTLRLDGLHGIGRGDID 662

RESULT 36

5180810-1

Patent No. 5180810

APPLICANT: Gomi, Hideyuki; Hozumi, Tatsunobu; Hattori, Shizuo;

Tagawa, Chiaki; Kishimoto, Fumitaka; Bjork, Lars

TITLE OF INVENTION: PROTEIN H CAPABLE OF BINDING TO ICG

NUMBER OF SEQUENCES: 4

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/07/376,641

FILING DATE: 07-JUL-1989

SEQ ID NO:1:

LENGTH: 376

5180810-1

Query Match 6.0%; Score 144; DB 6; Length 376;

Best Local Similarity 22.4%; Pred. No. 0.00016;

Matches 86; Conservative 49; Mismatches 117; Indels 132; Gaps 17;

Qy 68 QEEENVLDREFLK-----NELDNVRAQLSQDKKERSQVYIIDTLRDTLEERNATVVS 120

Db 49 QEEYKKLIEDNAKLVETTSLENEKTK-SENEKKN-----LDKLSKENG----- 95

Qy 121 LQALKAEMLCTLKKQKYLEQODETKQAQDEAGRLRSKMKTYEQIE----- 170

Db 96 -----GKLE-----KLELDYLLKLDHEKHEKQEQEQEERQKNOELERKYQREVERK 144

Qy 171 -----LLQSOLPEVEEMI-----RDMGVQSAVEQDLAVYCVSLKKEYEMLKEARKA 217

Db 145 YQEQLOKQOOL-ETEKQISASRKSLSRDLASRAKKDLAEHQKLEAEHQKLEKDKOI 203

Qy 218 SGEVADKLKRLDFSSRS-----KLQTVYSELDQAK-----LEL-KSA 253

Db 204 SDASRQGLSRDLASRAKKELAEANHQKLEAEHQKLEKDKOISDASRQGLSRDLASRAA 263

Qy 254 QKDL-----QSADEIMSLKKLTMLOETLNPVASETVDRVLVLESPAPVEYNLKLRRS 309

Db 264 KKELEANHOKLEAEARALKQAKQAEEL-----AKLR--- 296

Qy 310 FRDDIDLNATFDVTPPARPSSQHGYYEKLCLKESHSPIQDVPKKICKGPKRESQSLIG 369

Db 297 -----AGKASDSQOTDTRKGN-----KAYGKQQAQACGTRPNONKAPMKETKRQL- 342

Qy 370 GQSCAGEPDEELVGAPPIFYRNAI 393

Db 343 --PSTGET-----ANPFTAAAL 358

RESULT 37

PCT-US93-03077-3

Sequence 3, Application PC/TUS9303077

GENERAL INFORMATION:

APPLICANT: Board of Regents, The University of Texas System

APPLICANT: Gaynor, Richard B.

APPLICANT: Wu, Foon Kin

TITLE OF INVENTION: PROTEIN CELLULAR FACTOR USEFUL FOR

TITLE OF INVENTION: REGULATING GENE EXPRESSION

NUMBER OF SEQUENCES: 7

CORRESPONDENCE ADDRESS:

ADDRESSEE: Arnold White & Durkee

STREET: P.O. Box 4433

CITY: Houston

STATE: Texas

COUNTRY: USA  
ZIP: 77210  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: PCT/US93/03077  
FILING DATE: 19930331  
CLASSIFICATION:  
PRIOR APPLICATION DATA:  
APPLICATION NUMBER: US/07/862,025  
FILING DATE: April 2, 1992  
ATTORNEY/AGENT INFORMATION:  
NAME: Kammerer, Patricia A.  
REGISTRATION NUMBER: 29,775  
REFERENCE/DOCKET NUMBER: UFD270PCT  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: 713-787-1540  
TELEFAX: 713-749-2679  
TELEX:  
INFORMATION FOR SEQ ID NO: 3:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 414 amino acids  
TYPE: AMINO ACID  
STRANDEDNESS: unknown  
TOPOLOGY: unknown  
MOLECULE TYPE: peptide  
PCT-US93-03077-3

Query Match 6.0%; Score 144; DB 5; Length 414;  
Best Local Similarity 23.8%; Pred. No. 0.00018;

Matches 60; Conservative 47; Mismatches 77; Indels 68; Gaps 10;

QY 66 LAQEEENVLDREFLNELDNVRAQLSQDKERDSQYIIDLRLTL--EERNATVSLQ 122  
DB 5 LSEKEDVCKTVEFLNEKREARQLSLSEKALLEAFDMLKEMRVVESSISLTK 64  
QY 123 -----QALGRKEMLC---SLTKKQMKYLEQ-----QODETKQAOEE 155  
DB 65 DEFTQRIAEAEKKVQLACKERDAKKEIKIKELATRLNSSETADLLKEDEQIRGLME 124  
QY 156 AG-----RLRSKMKTMEOIELLOSOLPEVEEMIRMGV---GOSAVEQL 197  
DB 125 EGETLSQOQLNSNIITKLAKDKENEMVAKLKKVKELEEBELQHLKQVLDGKEVE-- 182  
QY 198 AVYCVSLAKKEYENLKEARKASGEVADKLRLKDLFSSRSKLQTVYSELDAQLELKSQKDL 257  
DB 183 -----KQRENIKRL-----NSMVERQEKDL---GRLOVDMDELEKN---RSIQAL 224  
QY 258 QSADEKINSJLK 269  
DB 225 DSAKKELTJDLHK 236

RESULT 38

US-08-533-306A-6

; Sequence 6, Application US/08533306A

; Patent No. 5837457

; GENERAL INFORMATION:

; APPLICANT: Liu, Pu

; APPLICANT: Collins, Francis S.

; APPLICANT: Sticiliano, Michael J.

; APPLICANT: Claxton, David

; TITLE OF INVENTION: Markers for Detection of Chromosome 16

; TITLE OF INVENTION: Rearrangements

; NUMBER OF SEQUENCES: 14

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Harness, Dickey & Pierce, P.L.C.

; STREET: P.O. Box 828

; CITY: Bloomfield Hills

STATE: MI  
COUNTRY: USA  
ZIP: 48303  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: PatentIn Release #1.0, Version #1.25  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/533,306A  
FILING DATE: September 25, 1995  
CLASSIFICATION: 435  
ATTORNEY/AGENT INFORMATION:  
NAME: Smith, Deann F.  
REGISTRATION NUMBER: 36683  
REFERENCE/DOCKET NUMBER: 2115-00869COB  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (810) 641-1600  
TELEFAX: (810) 641-0270  
INFORMATION FOR SEQ ID NO: 6:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 816 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-533-306A-6

Query Match 6.0%; Score 142.5; DB 2; Length 816;  
Best Local Similarity 22.6%; Pred. No. 0.00063;

Matches 76; Conservative 76; Mismatches 144; Indels 41; Gaps 15;

QY 61 KLFEDLAQEEENVLD--REFLNELD---NVRASQDKERDSQYIIDLRLTL----- 112  
DB 183 LKLDVASSUSSQDQVQLQETTRQKLNSTKLRLQEEENSLQ---DQLEDEMAKON 239  
QY 113 -ERNATVSLQALIGRAEM--LCSTL-----KKQKYLEQOODETKQAOEAGRL 159  
DB 240 LEHISTSLTNQLSDSKKQLDFASTYEALECGKKRRQKEIENLTQYEEKAAYDLKLT 299  
QY 160 RSKMKTMEOIELLOSOLPEVEEMIRMGVGSADVQLAVYCVSLAKKEYENLKEARKASG 219  
DB 300 KNRLQ--QELDDLTV--DLNQRLVSNLEKKQKFPQLLAEEKNISSKYAD--ERDRARA 354  
QY 220 EVADKLRLKDLFSSRSKLQTVYS--ELDOAKLELKSQKDLQSA---DDEIMSKKRLTM 273  
DB 355 EAREKETKALSLARALEBEAELEELRTNKKMLKAEMEDLVSSKDYGKRVHELESKRA 414  
QY 274 LQETL-NLPPVASETVDRVLESAP--VEVNLKLRPSFRDIDLNATFDVDTPPAPS 330  
DB 415 LETQMEMKQQLBELDELDQASDARKLRLEVNNQALKGF--ERDLQARDEQNEKKRQL 472  
QY 331 SSQ-HGYEKLCLEKSHSPIQDVPKKICKPKRESOL 366  
DB 473 QRQLHEYETELDEDERNRALAAAKKRLRGLDKLDEL 509

RESULT 39

US-08-742-923A-6

; Sequence 6, Application US/08742923A

; Patent No. 5869611

; GENERAL INFORMATION:

; APPLICANT: Liu, Pu

; APPLICANT: Collins, Francis S.

; APPLICANT: Sticiliano, Michael J.

; APPLICANT: Claxton, David

; TITLE OF INVENTION: Markers for Detection of Chromosome 16

; TITLE OF INVENTION: Rearrangements

; NUMBER OF SEQUENCES: 14

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: Harness, Dickey & Pierce, P.L.C.

; STREET: P.O. Box 828

; CITY: Bloomfield Hills

```
STATE: MI
COUNTRY: USA
ZIP: 48303
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/742,923A
FILING DATE: NO. 5869611ember 1, 1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Smith, Dean F.
REGISTRATION NUMBER: 36683
REFERENCE/DOCKET NUMBER: 2115-00869DVC
TELECOMMUNICATION INFORMATION:
TELEPHONE: (810) 641-1600
TELEFAX: (810) 641-0270
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 816 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-742-923A-6
```

```
Query Match 6.0%; Score 142.5; DB 2; Length 816;
Best Local Similarity 22.6%; Pred. No. 0.00063;
Matches 76; Conservative 76; Mismatches 144; Indels 41; Gaps 15;
```

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QY 61 KLFEDLAQEEENVLD--REFPKNELD---NVRAGLSQDKKRRSOYIIDLRPTLE---- 112
DB 183 KLAQVASTSSQLODTELLQEEETROKLVNSTRKRLQEEERNISQ---DQDEMEKON 239
QY 113 -ERNATVVSLOQALGKAEM--LCSTL-----KKQKYLEQOODETRKQAEAGRL 159
DB 240 LEHISTLNTIQLSDSKKRLQDFASTVEALEEGKKRFQKEIENLTQYEEKAAYDKLEKT 299
QY 160 RSKMKTEQIELLIQSOLPEVEEKIRDMGVGQSAVEOLAVYCVSLKKEYENLKERRASG 219
DB 300 KNRIQ--QELDDLVV--DLNQROLVSNLEKKQRKFDLLAEKNISSKYAD--ERDRAEA 354
QY 220 EVADKRLKDLFSSRSKLOTAYS--ELDOAKLELSAQKDLQSA---DKELMSLKKKLTW 273
DB 355 EAREKETKALSLARALEALEAKELERTKMKLKAEMEDLVSSKDDYGVKNVHELEKSKRA 414
QY 274 LQETL-NLPVASETVDRVLESFAP--VEVNLKLRPSFRDDIDLNATFDVDTPPARPS 330
DB 415 LEQMEEMKTLQLELEDELDQASEDAKRLRLVNMQALKGQF--ERDQARDEQNEKKRQL 472
QY 331 SSQ-HGYEKLCLKESHPIDVPKTKCGPKRESOL 366
DB 473 QROLHEYELEDEDERNERALAAAKKLEGLDLDEL 509
```

## RESULT 40

```
US-08-533-306A-4
Sequence 4, Application US/08533306A
Patent No. 5837457
GENERAL INFORMATION:
APPLICANT: Liu, Pu
APPLICANT: Collins, Francis S.
APPLICANT: Siciliano, Michael J.
APPLICANT: Claxton, David
TITLE OF INVENTION: Markers for Detection of Chromosome 16
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Harness, Dickey & Pierce, P.L.C.
STREET: P.O. Box 828
CITY: Bloomfield Hills
```

```
STATE: MI
COUNTRY: USA
ZIP: 48303
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/533,306A
FILING DATE: September 25, 1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Smith, Dean F.
REGISTRATION NUMBER: 36683
REFERENCE/DOCKET NUMBER: 2115-00869COB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (810) 641-1600
TELEFAX: (810) 641-0270
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 885 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-533-306A-4
```

```
Query Match 6.0%; Score 142.5; DB 2; Length 885;
Best Local Similarity 22.6%; Pred. No. 0.0007;
Matches 76; Conservative 76; Mismatches 144; Indels 41; Gaps 15;
```

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QY 61 KLFEDLAQEEENVLD--REFPKNELD---NVRAGLSQDKKRRSOYIIDLRPTLE---- 112
DB 252 KLAQVASTSSQLODTELLQEEETROKLVNSTRKRLQEEERNISQ---DQDEMEKON 308
QY 113 -ERNATVVSLOQALGKAEM--LCSTL-----KKQKYLEQOODETRKQAEAGRL 159
DB 309 LEHISTLNTIQLSDSKKRLQDFASTVEALEEGKKRFQKEIENLTQYEEKAAYDKLEKT 368
QY 160 RSKMKTEQIELLIQSOLPEVEEKIRDMGVGQSAVEOLAVYCVSLKKEYENLKERRASG 219
DB 369 KNRIQ--QELDDLVV--DLNQROLVSNLEKKQRKFDLLAEKNISSKYAD--ERDRAEA 423
QY 220 EVADKRLKDLFSSRSKLOTAYS--ELDOAKLELSAQKDLQSA---DKELMSLKKKLTW 273
DB 424 EAREKETKALSLARALEALEAKELERTKMKLKAEMEDLVSSKDDYGVKNVHELEKSKRA 483
QY 274 LQETL-NLPVASETVDRVLESFAP--VEVNLKLRPSFRDDIDLNATFDVDTPPARPS 330
DB 484 LEQMEEMKTLQLELEDELDQASEDAKRLRLVNMQALKGQF--ERDQARDEQNEKKRQL 541
QY 331 SSQ-HGYEKLCLKESHPIDVPKTKCGPKRESOL 366
DB 542 QROLHEYELEDEDERNERALAAAKKLEGLDLDEL 578
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Search completed: September 4, 2002, 16:10:49  
Job time: 7588 sec

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